

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/542,735
Source:	P.CT.
Date Processed by STIC:	07/28/2005

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER

VERSION-4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND

TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
 U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/542, 735	
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1Wrapped Nucleics Wrapped Aminos		
2Invalid Line Length	Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5Variable Length	Sequence(s)contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped	
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
(NEW ۱۰ مرزع)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11Use of <220>	Sequence(s)missing the <220> "Feature" and associated numeric identifiers and responses Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
"oug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid	



PCT

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/542,735**DATE: 07/28/2005

TIME: 15:43:48

Input Set : D:\LNK-002 Sequence Listing.txt
Output Set: N:\CRF4\07282005\J542735.raw

```
3 <110> APPLICANT: Biomay Produktions- und Handels-Aktiengesellschaft
      5 <120> TITLE OF INVENTION: Process for the preparation of hypoallergenic mosaic
              antigens
      8 <130> FILE REFERENCE: mosaic
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/542,735
                                                                  Does Not Comply
C--> 11 <141> CURRENT FILING DATE: 2005-07-20
                                                                  Corrected Diskette Needed
     13 <160> NUMBER OF SEQ ID NOS: 12
     15 <170> SOFTWARE: PatentIn Ver. 2.1
     17 <210> SEQ ID NO: 1
     18 <211> LENGTH: 103
    19 <212> TYPE: PRT
    20 <213> ORGANISM: Artificial Sequence
     22 <220> FEATURE:
    23 <223> OTHER INFORMATION: Description of Artificial Sequence: rearranged
    24 polypeptide sequence
26 <400> SEQUENCE: 1
    27 Met Val Pro Lys Val Thr Phe Thr Val Glu Lys Gly Ser Asn Glu Lys
    28
                          5
        1
    30 His Leu Ala Val Leu Val Lys Tyr Glu Gly Asp Thr Met Ala Glu Val
    31
                     20
                                         25
    33 Glu Leu Phe Arg Phe Leu Thr Glu Lys Gly Met Lys Asn Val Phe Asp
                35
                                     40
    36 Asp Val Val Pro Glu Lys Tyr Thr Ile Gly Ala Thr Tyr Ala Pro Glu
                                 55
    39 Glu Arg Glu His Gly Ser Asp Glu Trp Val Ala Met Thr Lys Gly Glu
                             70
                                                 75
    42 Gly Gly Val Trp Thr Phe Asp Ser Glu Glu Pro Leu Gln Gly Pro Phe
    45 Asn His His His His His
    46
                    100
    49 <210> SEQ ID NO: 2
    50 <211> LENGTH: 309
    51 <212> TYPE: DNA
    52 <213> ORGANISM: Artificial Sequence
    54 <220> FEATURE:
    55 <223> OTHER INFORMATION: Description of Artificial Sequence: rearranged
    56
             (nucleotide sequence
    58 <400> SEQUENCE: 2
    59 atggtcccga aggtgacgtt cacggtggag aaggggtcca acgagaagca cctggcggtg 60
    60 ctggtgaagt acgaggggga caccatggcg gaggtggagc tcttccggtt cctcaccgag 120
    61 aagggcatga agaacgtctt cgacgacgtc gtcccagaga agtacaccat tggggccacc 180
    62 tacgcgccag aagagcggga gcacggctcc gacgagtggg tcgccatgac caagggggag 240
    63 ggcggcgtgt ggacgttcga cagcgaggag ccgctccagg ggcccttcaa ccaccaccac 300
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> See green summary sheet
               Input Set : D:\LNK-002 Sequence Listing.txt
               Output Set: N:\CRF4\07282005\J542735.raw
64 caccaccac
67 <210> SEQ ID NO: 3
68 <211> LENGTH: 34
69 <212> TYPE: PRT
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Description of Artificial Sequence: polypeptide
75 <400> SEQUENCE: 3
76 Val Pro Lys Val Thr Phe Thr Val Glu Lys Gly Ser Asn Glu Lys His
                                       10
79 Leu Ala Val Leu Val Lys Tyr Glu Gly Asp Thr Met Ala Glu Val Glu
                                   25
80
82 Leu Cys
                                                          J Same Error
86 <210> SEQ ID NO: 4
87 <211> LENGTH: 33
88 <212> TYPE: PRT
89 <213> ORGANISM: Artificial Sequence
91 <220> FEATURE:
92 <223> OTHER INFORMATION: (Description of Artificial Sequence: polypeptide
94 <400> SEQUENCE: 4
95 Arg Glu His Gly Ser Asp Glu Trp Val Ala Met Thr Lys Gly Glu Gly
                    5
                                       10
98 Gly Val Trp Thr Phe Asp Ser Glu Glu Pro Leu Gln Gly Pro Phe Asn
               20
99
                                   25
101 Cys
                                                                      -7 Same Error
105 <210> SEQ ID NO: 5
106 <211> LENGTH: 32
107 <212> TYPE: PRT
108 <213> ORGANISM: Artificial Sequence
110 <220> FEATURE:
111 <223> OTHER INFORMATION: (Description of Artificial Sequence: polypeptide
113 <400> SEQUENCE: 5
114 Cys Phe Arg Phe Leu Thr Glu Lys Gly Met Lys Asn Val Phe Asp Asp
                    - 5
117 Val Val Pro Glu Lys Tyr Thr Ile Gly Ala Thr Tyr Ala Pro Glu Glu
                                    25
124 <210> SEQ ID NO: 6
125 <211> LENGTH: 34
126 <212> TYPE: DNA
127 <213> ORGANISM: Artificial Sequence
129 <220> FEATURE:
130 <223> OTHER INFORMATION: Description of Artificial Sequence: primer
132 <400> SEQUENCE: 6
133 qgatttccat atggtcccga aggtgacgtt cacg
                                                                     34
136 <210> SEQ ID NO: 7
137 <211> LENGTH: 36
138 <212> TYPE: DNA
139 <213> ORGANISM: Artificial Sequence
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/542,735

141 <220> FEATURE:

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/542,735**DATE: 07/28/2005

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Input Set : D:\LNK-002 Sequence Listing.txt
Output Set: N:\CRF4\07282005\J542735.raw

142 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 144 <400> SEQUENCE: 7 145 ggtgaggaac cggaagagct ccacctccgc catggt 36 148 <210> SEQ ID NO: 8 149 <211> LENGTH: 36 150 <212> TYPE: DNA 151 <213> ORGANISM: Artificial Sequence 153 <220> FEATURE: 154 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 156 <400> SEQUENCE: 8 157 gcggaggtgg agctcttccg gttcctcacc gagaag 36 160 <210> SEQ ID NO: 9 161 <211> LENGTH: 36 162 <212> TYPE: DNA 163 <213> ORGANISM: Artificial Sequence 165 <220> FEATURE: 166 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 168 <400> SEQUENCE: 9 36 169 ggagccgtgc tcccgctctt ctggcgcgta ggtggc 172 <210> SEQ ID NO: 10 173 <211> LENGTH: 36 174 <212> TYPE: DNA 175 <213> ORGANISM: Artificial Sequence 177 <220> FEATURE: 178 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 180 <400> SEQUENCE: 10 36 181 tacgcgccag aagagcggga gcacggctcc gacgag 184 <210> SEQ ID NO: 11 185 <211> LENGTH: 51 186 <212> TYPE: DNA 187 <213> ORGANISM: Artificial Sequence 189 <220> FEATURE: 190 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 192 <400> SEQUENCE: 11 193 cgcgaattet cagtggtggt ggtggtggtg gttgaaggge ceetggageg g 51 196 <210> SEO ID NO: 12 197 <211> LENGTH: 51 198 <212> TYPE: DNA 199 <213> ORGANISM: Artificial Sequence 201 <220> FEATURE: 202 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 204 <400> SEQUENCE: 12 205 cgcgaattct cagtggtggt ggtggtggtg ctcttctggc gcgtaggtgg c 51

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VERIFICATION SUMMARY

DATE: 07/28/2005

PATENT APPLICATION: US/10/542,735

TIME: 15:43:50

Input Set : D:\LNK-002 Sequence Listing.txt Output Set: N:\CRF4\07282005\J542735.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date